

Table E1: Sequences of PCR Primers

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3 4 5	<u>Gene</u>	<u>Sequence</u>	Product Size (bp)
6 7 8	CD68 sense CD68 antisense	AGATTCGAGTCATGTACACAACCCA [SEQ GGTGCTTGGAGATCTCGAAG [SEQ ID NO:2]	ID NO:1] 279
9 10 11	$P_{2Y1}R$ sense $P_{2Y1}R$ antisense	TGTGGTGTACCCCCTCAAGTCCC [SEQ ID NO ATCCGTAACAGCCCAGAATCAGCA [SEQ ID	
12 13 14	$P_{2Y2}R$ sense $P_{2Y2}R$ antisense	CCAGGCCCCCGTGCTCTACTTTG (SEQ ID NO CATGTTGATGGCGTTGAGGGTGTG[SEQ ID	D:5] 367 NO:6]
15 16 17	CXCR4 sense CXCR4 antisense	TTCTACCCCAATGACTTGTG [SEQ ID NO:7] ATGTAGTAAGGCAGCCAACA [SEQ ID NO:8]	206
18 19	MIP-1α sense MIP-1α antisense	ACCATGGCTCTCTGCAACCA [SEQ ID NO:9] TTAAGAAGAGTCCCACAGTG[SEQ IDNO:10]	393
20 21 22	MIP-1β sense MIP-1β antisense	CCTGCTGCTTTTCTTACACC [SEQ ID NO:11] CACCTAATACAATAACACGGC [SEQ ID NO:1	336
23 24 25	MCP-1 sense MCP-1 antisense	ATAGCAGCCACCTTCATTCC [SEQ ID NO:13] TTCCCCAAGTCTCTGTATCT [SEQ ID NO:14]	466
26 27 28	IL-1β sense IL-1β antisense	AAAAGCTTGGTGATGTCTGG [SEQ ID NO:15] TTTCAACACGCAGGACAGG [SEQ ID NO:16]	179
29 30 31	IL-2 sense IL-2 antisense	ATGGTTGCTGTCTCATCAGC [SEQ ID NO:17] CTGGAGCATTTACTGCTGGA [SEQ ID NO:18]	
32 33 34	IL-3 sense IL-3 antisense	ATGAGCCGCCTGCCCGTCCTG [SEQ ID NO:1 AAGATCGCGAGGCTCAAAGTCGTCTGTTC	
35 36 37	IL-4 sense IL-4 antisense	GACACAAGTGCAATATCACC (SEQ ID NO:21 AAGTTTTCCAACGTACTCTG (SEQ ID NO:22)	l 337
38 39 40	IL-5 sense IL-5 antisense	GAGGATGCTTCTGCATTTGAGTTTG (SEQ II GTCAATGTATTTCTTTATTAAGGACAAG (S	
41 42 43 44	IL-6 sense IL-6 antisense	GTGTGAAAGCAGCAAAGAGGC (SEQ ID NO: CTGGAGGTACTCTAGGTATAC (SEQ ID NO: 2	

## Table E1: Sequences of PCR Primers (continued)

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3 4 5	Gene	Sequence Product Si (bp)	<u>ze</u>
6 7 8	IL-7 sense IL-7 antisense	TGTTGAACTGCACTGGCCAG [SEQ ID NO:27] GCAACTGATACCTTACATGG [SEQ ID NO:28]	484
9 10 11	IL-8 sense IL-8 antisense	ATGACTTCCAAGCTGGCCGTG [SEQ ID NO:29] TATGAATTCTCAGCCCTCTTCAAAA [SEQ ID NO:30]	<b>30</b> 1
12 13 14	IL-9 sense IL-9 antisense	ATGCTTCTGGCCATGGTCCT [SEQ ID NO:31] TATCTTGCCTCTCATCCCTC [SEQ ID NO:32]	375
15 16 17	IL-10 sense IL-10 antisense	AGATCTCCGAGATGCCTTCAGCAGA [SEQ ID NO:33] CCTTGATGTCTGGGTCTTGGTTCTC [SEQ ID NO:34]	194
18 19 20	IL-11 sense IL-11 antisense	ACTGCTGCTGCAAGACTCGGCTGTGA [SEQ ID NO:3 ATGGGGAAGAGCCAGGGCAGAAGTCTGT [SEQ ID NO:	-
21 22 23	IL-12 sense IL-12 antisense	TCACAAAGGAGGCGAGGTTCTAAGC [SEQ ID NO:37] CCTCTGCTGCTTTTGACACTGAATG [SEQ ID NO:38]	213
24 25 26	IL-13 sense IL-13 antisense	ACCCAGAACCAGAAGGCTCCG [SEQ ID NO:39] TCAGTTGAACCGTCCCTGGCG [SEQ ID NO:40]	198
27 28 29	IL-15 sense IL-15 antisense	AAACCCCCTGCCATAGCCAACTCTT [SEQ ID NO:41] CTTCTGTTTTAGGGAGCCCTGCACT [SEQ ID NO:42]	202
30 31	TNF- $\alpha$ sense TNF- $\alpha$ antisense	CAAAGTAGACCTGCCCAGAC [SEQ ID NO:43] GACCTCTCTCAATCAGCCC [SEQ ID NO:44]	490
32 33 34	NF-M sense NF-M antisense	TGGGAAATGGCTCGTCATTT [SEQ ID NO:45] CTTCATGGAAGCGGCCAATT [SEQ ID NO:46]	333
35 36 37	MBP sense MBP antisense	ACACGGGCATCCTTGACTCCATCGG [SEQ ID NO:47] TCCGGAACCAGGTGGGTTTTCAGCG [SEQ ID NO:48]	510
38 39 40	GFAP sense GFAP antisense	GCAGAGATGATGGAGCTCAATGACC [SEQ ID NO:49] GTTTCATCCTGGAGCTTCTGCCTCA [SEQ ID NO:50]	266
41 42 43 44	B7-2 sense B7-2 antisense	CTCTTTGTGATGGCCTTCCTG [SEQ ID NO:51] CTTAGGTTCTGGGTAACCGTG [SEQ ID NO:52]	464